



ENTERED

PCT09

#9

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/674,752

DATE: 02/14/2003

TIME: 13:19:07

Input Set : A:\EP.txt

Output Set: N:\CRF4\02142003\I674752.raw

3 <110> APPLICANT: Voorberg, Johannes  
 5 <120> TITLE OF INVENTION: Method For Diagnosis and Treatment of Haemophilia A Patients  
 With An  
 6 Inhibitor  
 8 <130> FILE REFERENCE: Sequence Nos 1-59 for 294-86 PCT/US  
 10 <140> CURRENT APPLICATION NUMBER: 09/674,752  
 11 <141> CURRENT FILING DATE: 2000-12-29  
 13 <150> PRIOR APPLICATION NUMBER: PCT/NL99/00285  
 14 <151> PRIOR FILING DATE: 1999-05-07  
 16 <150> PRIOR APPLICATION NUMBER: EP 98201543.0  
 17 <151> PRIOR FILING DATE: 1998-05-08  
 19 <160> NUMBER OF SEQ ID NOS: 59  
 21 <170> SOFTWARE: PatentIn version 3.1  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 24  
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 26 <213> ORGANISM: Homo sapiens  
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 35 <213> ORGANISM: Homo sapiens  
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 61 <212> TYPE: DNA  
 62 <213> ORGANISM: Homo sapiens  
 64 <400> SEQUENCE: 5  
 65 gaggtgcagc tgggtggagtc tgg 23  
 68 <210> SEQ ID NO: 6

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83 gaggtacagc tgcagcagtc tgc 23
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89 <213> ORGANISM: Homo sapiens
91 <400> SEQUENCE: 8
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96 <211> LENGTH: 36
97 <212> TYPE: DNA
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132 <211> LENGTH: 28
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160 <212> TYPE: DNA
161 <213> ORGANISM: Homo sapiens
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188 <213> ORGANISM: Homo sapiens
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195 cctggacaag ggcttgagtg gatgggaggg atcatcccta tctttggttc aacaaagtac 180
197 gcacagaagt tccagggcag agtcacgatg accgcggacg gatccacgag tacagcctac 240
199 atggaactga acagcctgag atctgaggac acggccatat attactgtgc gcgacaacag 300
201 aacggcggct ggtacgaagg accgttgctt gagccgaggc ctgatgctct tgatatctgg 360
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216 cctggacaag ggcttgagtg gatgggaggg atcatcccta tctttggtac agcaaaactac 180
218 gcacagaagt tccagggcag agtcacgatt accgcggacg aatccacgag cacagcctac 240

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231 tcctgcatgg cttctgggta cccctttacc agctatgata tcagttgggt gcgacaggcc      120
233 cctggacaag ggcttgagtg gatgggatgg atcagcattt atagtggtaa cacagactat      180
235 gcacagaagt tccagggcag agtcaccatg acgacagaca catccaggag aacagcctac      240
237 atggagctga ggagcctgag atctgacgac acggccgtct attattgtgc gagagatggg      300
239 ggggggggtg cctatgaaga tgtttggagt ggtgagtacc ccgaatacta cgctatggac      360
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254 cctggacaag ggcttgagtg gatgggatgg atcagcgctt acaatggtaa cacaaactat      180
256 gcacagaagc tccagggcag agtcaccatg accacagaca catccacgag cacagcctac      240
258 atggagctga ggagcctgag atctgacgac acggccgtgt attactgtgc gagaga      296
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263 <212> TYPE: PRT
264 <213> ORGANISM: Homo sapiens
266 <400> SEQUENCE: 23
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272 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Asp Thr Phe Asn Ser Phe
273          20          25          30
276 Pro Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
277          35          40          45
280 Gly Gly Ile Ile Pro Ile Phe Gly Ser Thr Lys Tyr Ala Gln Lys Phe
281          50          55          60
284 Gln Gly Arg Val Thr Met Thr Ala Asp Gly Ser Thr Ser Thr Ala Tyr
285 65          70          75          80
288 Met Glu Leu Asn Ser Leu Arg Ser Glu Asp Thr Ala Ile Tyr Tyr Cys
289          85          90          95
292 Ala Arg Gln Gln Asn Gly Gly Trp Tyr Glu Gly Pro Leu Leu Glu Pro
293          100         105         110
296 Arg Pro Asp Ala Leu Asp Ile Trp Gly Gln Gly Thr Met Val Thr Val
297          115         120         125
300 Ser Ser
301          130
304 <210> SEQ ID NO: 24
305 <211> LENGTH: 98
306 <212> TYPE: PRT

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307 <213> ORGANISM: Homo sapiens
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312 1 5 10 15
315 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser Tyr
316 20 25 30
319 Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
320 35 40 45
323 Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
324 50 55 60
327 Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
328 65 70 75 80
331 Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
332 85 90 95
335 Ala Arg
339 <210> SEQ ID NO: 25
340 <211> LENGTH: 132
341 <212> TYPE: PRT
342 <213> ORGANISM: Homo sapiens
344 <400> SEQUENCE: 25
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347 1 5 10 15
350 Ser Met Lys Val Ser Cys Met Ala Ser Gly Tyr Pro Phe Thr Ser Tyr
351 20 25 30
354 Asp Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
355 35 40 45
358 Gly Trp Ile Ser Ile Tyr Ser Gly Asn Thr Asp Tyr Ala Gln Lys Phe
359 50 55 60
362 Gln Gly Arg Val Thr Met Thr Thr Asp Thr Ser Arg Arg Thr Ala Tyr
363 65 70 75 80
366 Met Glu Leu Arg Ser Leu Arg Ser Asp Asp Thr Ala Val Tyr Tyr Cys
367 85 90 95
370 Ala Arg Asp Gly Gly Gly Ala Tyr Glu Asp Val Trp Ser Gly Glu
371 100 105 110
374 Tyr Pro Glu Tyr Tyr Ala Met Asp Val Trp Gly Gln Gly Thr Thr Val
375 115 120 125
378 Thr Val Ser Ser
379 130
382 <210> SEQ ID NO: 26
383 <211> LENGTH: 98
384 <212> TYPE: PRT
385 <213> ORGANISM: Homo sapiens
387 <400> SEQUENCE: 26
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390 1 5 10 15
393 Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr
394 20 25 30
397 Gly Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
398 35 40 45

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/674,752

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Input Set : A:\EP.txt

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